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Figure 1

NCE4	MRSSPLLRSAVVAALPVLALAADGKSTRYWDCCKPSCGWAK	21
STCE1	********** * MRSSPVLRTALAAALPLAALAADGKSTRYWDCCKPSCSWPG	20
NCE5	* ****** *** MQLPLTTLLTLLPALAAAQSGSGRTTRYWDCCKPSCAWPG Signal peptide → Catalytic domain →	23
	Signal peptide → Catalytic domain →	
NCE4	KAPVNQPVFSCNANFQRLTDF-DAKSGCEPGGVAYSCADQTPWAVNDDFA	70
STCE1	** ***** * **** * *** ** ** ****** * KASVNQPVFACSANFQRISDP-NVKSGCD-GGSAYACADQTPWAVNDNFS	68
NCF5	* ** * * * * *** ** * *** * KGPA—PVRTCDRWDNPLFDGGNTRSGCDAGGGAYMCSDQSPWAVSDDLA—	71
NOES	RGFAFVRTODR#DNFLFDGGNTRSGCDAGGATMCSDGSFNAVSDDLA-	/ 1
NCE4	FGFAATS AGSNEAGWCCACYELTFTSGPVAGKKMVVQSTSTGGDLGSNH ******** * *** *** ******************	120
STCE1	YGFAATS I SGGNEASWCCGCYELTFTSGPVAGKTMVVQSTSTGGDLGTNH	118
NCE5	** ** * * ** *** ********** * ** ******	121
NCE4	FDLNIPGGGVGIFDGCTPQFGGLPGQRYGGISSRNECDRFPDALKPG *** ******* ****** * ***************	167
STCE1	FDLAMPGGGVG1FDGCSPQFGGLAGDRYGGVSSRSQCDSFPAALKPG	165
NCE5	** ******* * * * * **** * ** **** FDIAMPGGGVGIFNACTDQYGAPPNGWGQRYGGISQRHECDAFPEKLKPG	171

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Figure 2

			217
NCE4	CYWRFDWFKNADNPSFSFRQVQCPAELVARTGCRRNDDGNFPAVQ1PSSS		
	solololololololololok w sololololok solololololok	skskstotoksk sksk	
STCE1	CYWRFDWFKNADNPTFTFRQVQCPSELVARTGCR	RNDDGNFPVFTPPSGG	215
	stateletatek stateletek statek state state	*	
NCE5	CYWRFDWFLNADNPSVNWRQVSCPAE I VAKSGCS	R	206
11020		 Linker→	200
		LITIKEI -	
NCE4	TSSPVGQPTSTSTTSTSTTSSPPVQPTTPS	GCTAERWA	255
	** * ***** * *	** * **	
STCE1	QSSSSSSSSSAKPTSTSTSTTSTKATSTTSTASS	OTSSSTGGGCAAQRWA	265
NCE5			!
NOLO		0.0.0	
		CBD→	
:			
NCE4	CQCGGNGWSGCTTCVAGSTCTKINDWYHQCL	286	
	kololotek ak atololotekek ak ak ak akololotekek		
STCE1	CQCGGIGFSGCTTCVSGTTCNKQNDWYSQCL	295	
	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		
NOTE			
NCE5			